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O I P E

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/830,703

DATE: 08/30/2001
TIME: 06:40:43

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\08302001\1830703.raw

4 <110> APPLICANT: Biofrontera Pharmaceuticals AG
5 Lubbert, Hermann
7 <120> TITLE OF INVENTION: TRANSGENIC ANIMAL MODEL FOR
8 NEURODEGENERATIVE DISEASES
10 <130> FILE REFERENCE: STERN1.001APC
12 <140> CURRENT APPLICATION NUMBER: 09/830,703
13 <141> CURRENT FILING DATE: 2001-04-26
15 <150> PRIOR APPLICATION NUMBER: EP 99116766.9
16 <151> PRIOR FILING DATE: 1999-08-30
18 <160> NUMBER OF SEQ ID NOS: 34
20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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23 <211> LENGTH: 3255
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25 <213> ORGANISM: Mus musculus
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30 cggtgaccat gatagtgttt gtcagggttca actccagcta tggcttccca gtggaggtcg 180
31 attctgacac cagcatctt cagctcaagg aagtgggtgc taagcgacag ggggttccag 240
32 ctgaccagct gctgtgttattttccggga aggagcttcc gaatcacctg acggttcaaa 300
33 actgtgacact ggaacaacag agtattgtac acatagtaca gagaccacgg aggagaagtc 360
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35 agtccaggag cttgacacga gtggacactga gcagccatac cctggccgtg gactctgtgg 480
36 ggctggcggt cattctggac acagacagta agagggattc agaagcagcc agagggtccag 540
37 ttaaacccac ctacaacacg ttttcatct actgcaaagg cccctgccc aaggtcac 600
38 ctggaaagct ccgagttcag tttggcacct gcaaacaagc aaccctcacc ttggccagg 660
39 gcccacatctt cttggacat gtcttaattt caaacggat gagttgtgag tgccagtc 720
40 cagactgccc tggaaaccaga gctgaatttt tctttaatg tggagcacac ccaacccat 780
41 acaaggacac gtcggtagct ttgaacactga tcaccagcaa cagggcgcac atccctgca 840
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44 acgatgctca acttggctac tccctggcggt gtgttagctgg ctgtcccaac tccctgatta 1020
45 aagagctcca tcacttcagg atccttggag aagagcagta cactaggtac cagcagttat 1080
46 gggccgagga atgcgtgtc caaatgggag gtgtgtgtc ccccccgtcct ggctgtggag 1140
47 ctggactgtc acctgaacag gccagaggg aagtccatcg cgaagggggc aacggcctgg 1200
48 gctgcgggtt tgtttctgc cggactgtc aggaagcata ccatgaaggg gattgcact 1260
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57 ctctggctca ctcgaatgtc agagcttaga tgaggggtgca cctttgggtt ggattctgt 1800
58 qaagccatga gtgaggtggg aagtgtttc cagggttgtt gccacccct gggtaagta 1860

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61 cctatctta caaatggtgc tcatggagatt acaaccctc tggactaa tcagcttatac 2040
62 aaccaagtga gaaccttagga aagctaattt gatggcagac tgcttaaattc gcagggagga 2100
63 ctcagaagcc aaacctactt ccgttcgtt cattatctgc aacttttagaa agaaatgatc 2160
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66 ttaagagaca atcccatgt gctgcgccaa tgctaaacac agggtaaga aacacaatgt 2340
67 ttatagaagg agcatccctc gaccatctga atgagagttt gctgaccctt ttccaccaca 2400
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69 ccacccagtg gtcctacaga cagggcaata cacacacacc aagatagcc tcagatcaac 2520
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71 tttttgtttt ttgctttttt tttttttttt tttgggtggg gtggggctac caaacttgag 2640
72 gcctagagct aaaaatcata tagaaatgtt gttatctgtt ggtgtgagga aaggccagct 2700
73 ggccctaagtt cacacttttgc tcccaactggc cctagactcc acccagccag ctcccaaaat 2760
74 gaaaagacca cctgtcaagc agcagtcaagg agtctgtatgt cacccatcac tattttttt 2820
75 ccatcattgt gcttgcctct gcctccttcc acaccgtgt gacgtaatcg cattgggaag 2880
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78 attcttaaaa taacccaaaaa gacaggcattt cacagtgtgtt gacatgaat cacgcctgc 3060
79 attgtgtgag tggatgtt gggataaaag tggatgtcag aagatggaa atcaaaccctc 3120
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92 tggatgtgtt tggatgttttca aactccagct atggcttccc agtggagggtt gattctgaca 180
93 ccagcatctt gcagctcaag gaagtgggg ttaagcgaca ggggggttcca gctgaccagc 240
94 tgcgtgtat ttttgcggg aaggagcttgc cgaatcacct gacggttcaa aactgtgacc 300
95 tggacaaca gatgttttgc cacatgttgc agagaccacg gaggagaagt catgaaacaa 360
96 atgcatctgg agggggacgaa ccccaagagca cctcagaggg ctccatatgg gatccagga 420
97 gcttgcacacg agtggacacgtt agcagccata ccctggcggt ggactctgtt gggctggcg 480
98 tcattctggaa cacagacagt aagaggattt cagaagcagc cagagggttcca gcagttaaac 540
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100 agtcccgagt tcagttgttgc acctgcaaaac aagcaaccctt caccttggcc cagggcccat 660
101 cttgtgtggaa cgtatgttta attccaaacc ggtatgttgc tgatgttgc tctccagact 720
102 gcccgttggaa cagagctgaa ttttttttta aatgtggggc acacccaaacc tcagacaagg 780
103 acacgtcggtt agcttgcacca ctgatcacca gcaacaggcg cagcatccct tgcatacggt 840
104 gcacagatgtt cgtatctt cctctgtcat ctggcgccctc cgtgtggactt cggccctatc 900
105 tccactgaac cttgttctt aggactgttgc aataggtgtt cacccttcttac tgagaacaag 960
106 gcagcttctg gtcttgcgtt tcccttgcgtt ccaacggcg cattgactgtt acacccttca 1020
107 gtccttaccaa ccccaattacc tgggttgcgtt ctttaccgtt tagcttcttcc aagatgccta 1080
108 tttccacaca cagtttcttgc tcttccat ccccccattttt gtttatgttgc atgatgttgc 1140
109 accgcacccatc atgatgttgcgtt gtttgcgttgcata caagacttcc tggatcccc gcttgcgttgc 1200

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 111 tcttgggctg gtgagatggc tcagtggtt agagcacccg actgctctc cgaagtccag 1320
 112 agttcaaaat cccagcaacc acatggtggc tcacaaccat ccgtaacaag atctgactcc 1380
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 124 gttcaactcc agctatggct tcccagtgg agtcgattct gacaccagca tcttgcagct 180
 125 caaggaagtg gttgctaaac gacagggggt tccagctgac cagctgcgtg tgattttgc 240
 126 cgggaaggag cttccgaatc acctgacggt tcaaaactgt gacctggAAC aacagagtt 300
 127 ttttacacata gtacagagac cacggaggag aagtcatgaa acaaattgtat ctggagggg 360
 128 cgaaccccaag agcaccttag agggctccat atgggagttcc aggagcttga cacgagtgg 420
 129 cctgagcagc cataccctgc cggggactc tggggctg gcggcatttc tggacacaga 480
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 131 tttcatctac tgcaaaggcc cctgccacaa ggtccagctt gggaaagctcc gagttcagtg 600
 132 tggcacctgc aaacaagcaa ccctcacctt ggccagggtt ccattttgtt gggacgtgt 660
 133 ctttaatttcca aaccggatga gtggtgatgtt ccagtcttca gactgcctt gaaaccagac 720
 134 tgaatttttc tttaaatgtt gggcacaccc aacccatgtt aaggacacgtt cggtagctt 780
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 136 tatgcgcatg agttac 857
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 139 <211> LENGTH: 464
 140 <212> TYPE: PRT
 141 <213> ORGANISM: Mus musculus
 143 <400> SEQUENCE: 4
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 147 20 25 30
 148 Arg Gln Gly Val Pro Ala Asp Gln Leu Arg Val Ile Phe Ala Gly Lys
 149 35 40 45
 150 Glu Leu Pro Asn His Leu Thr Val Gln Asn Cys Asp Leu Glu Gln Gln
 151 50 55 60
 152 Ser Ile Val His Ile Val Gln Arg Pro Arg Arg Arg Ser His Glu Thr
 153 65 70 75 80
 154 Asn Ala Ser Gly Gly Asp Glu Pro Gln Ser Thr Ser Glu Gly Ser Ile
 155 85 90 95
 156 Trp Glu Ser Arg Ser Leu Thr Arg Val Asp Leu Ser Ser His Thr Leu
 157 100 105 110
 158 Pro Val Asp Ser Val Gly Leu Ala Val Ile Leu Asp Thr Asp Ser Lys
 159 115 120 125
 160 Arg Asp Ser Glu Ala Ala Arg Gly Pro Val Lys Pro Thr Tyr Asn Ser
 161 130 135 140
 162 Phe Phe Ile Tyr Cys Lys Gly Pro Cys His Lys Val Gln Pro Gly Lys

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163 145 150 155 160
 164 Leu Arg Val Gln Cys Gly Thr Cys Lys Gln Ala Thr Leu Thr Leu Ala
 165 165 170 175
 166 Gln Gly Pro Ser Cys Trp Asp Asp Val Leu Ile Pro Asn Arg Met Ser
 167 180 185 190
 168 Gly Glu Cys Gln Ser Pro Asp Cys Pro Gly Thr Arg Ala Glu Phe Phe
 169 195 200 205
 170 Phe Lys Cys Gly Ala His Pro Thr Ser Asp Lys Asp Thr Ser Val Ala
 171 210 215 220
 172 Leu Asn Leu Ile Thr Ser Asn Arg Arg Ser Ile Pro Cys Ile Ala Cys
 173 225 230 235 240
 174 Thr Asp Val Arg Ser Pro Val Leu Val Phe Gln Cys Asn His Arg His
 175 245 250 255
 176 Val Ile Cys Leu Asp Cys Phe His Leu Tyr Cys Val Thr Arg Leu Asn
 177 260 265 270
 178 Asp Arg Gln Phe Val His Asp Ala Gln Leu Gly Tyr Ser Leu Pro Cys
 179 275 280 285
 180 Val Ala Gly Cys Pro Asn Ser Leu Ile Lys Glu Leu His His Phe Arg
 181 290 295 300
 182 Ile Leu Gly Glu Glu Gln Tyr Thr Arg Tyr Gln Gln Tyr Gly Ala Glu
 183 305 310 315 320
 184 Glu Cys Val Leu Gln Met Gly Gly Val Leu Cys Pro Arg Pro Gly Cys
 185 325 330 335
 186 Gly Ala Gly Leu Leu Pro Glu Gln Gly Gln Arg Lys Val Thr Cys Glu
 187 340 345 350
 188 Gly Gly Asn Gly Leu Gly Cys Gly Phe Val Phe Cys Arg Asp Cys Lys
 189 355 360 365
 190 Glu Ala Tyr His Glu Gly Asp Cys Asp Ser Leu Leu Glu Pro Ser Gly
 191 370 375 380
 192 Ala Thr Ser Gln Ala Tyr Arg Val Asp Lys Arg Ala Ala Glu Gln Ala
 193 385 390 395 400
 194 Arg Trp Glu Glu Ala Ser Lys Glu Thr Ile Lys Lys Thr Thr Lys Pro
 195 405 410 415
 196 Cys Pro Arg Cys Asn Val Pro Ile Glu Lys Asn Gly Gly Cys Met His
 197 420 425 430
 198 Met Lys Cys Pro Gln Pro Gln Cys Lys Leu Glu Trp Cys Trp Asn Cys
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 207 <213> ORGANISM: Mus musculus
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 214 Arg Gln Gly Val Pro Ala Asp Gln Leu Arg Val Ile Phe Ala Gly Lys

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220	Asn	Ala	Ser	Gly	Gly	Asp	Glu	Pro	Gln	Ser	Thr	Ser	Glu	Gly	Ser	Ile
221															95	
222	Trp	Glu	Ser	Arg	Ser	Leu	Thr	Arg	Val	Asp	Leu	Ser	Ser	His	Thr	Leu
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224	Pro	Val	Asp	Ser	Val	Gly	Leu	Ala	Val	Ile	Leu	Asp	Thr	Asp	Ser	Lys
225														125		
226	Arg	Asp	Ser	Glu	Ala	Ala	Arg	Gly	Pro	Ala	Val	Lys	Pro	Thr	Tyr	Asn
227														140		
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229	145		150		155		160									
230	Lys	Leu	Arg	Val	Gln	Cys	Gly	Thr	Cys	Lys	Gln	Ala	Thr	Leu	Thr	Leu
231														175		
232	Ala	Gln	Gly	Pro	Ser	Cys	Trp	Asp	Asp	Val	Leu	Ile	Pro	Asn	Arg	Met
233														190		
234	Ser	Gly	Glu	Cys	Gln	Ser	Pro	Asp	Cys	Pro	Gly	Thr	Arg	Ala	Glu	Phe
235														205		
236	Phe	Phe	Lys	Cys	Gly	Ala	His	Pro	Thr	Ser	Asp	Lys	Asp	Thr	Ser	Val
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264	Trp	Glu	Ser	Arg	Ser	Leu	Thr	Arg	Val	Asp	Leu	Ser	Ser	His	Thr	Leu
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